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SEQUENCE LISTING

<110> Qian, Su
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Metzger, Joseph M.

<120> Agouti-related protein deficient cells, non-humkan transgenic animals and methods of selecting compounds which regulate energy metabolism

<130> 21033YP

<150> PCT/US03/20245
<151> 2003-06-27

<150> 60/393,391
<151> 2002-07-03

<160> 14

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        35                  40                  45
Gly Leu Lys Lys Thr Thr Ala Asp Arg Ala Glu Glu Val Leu Leu Gln
    50                  55                  60
Lys Ala Glu Ala Leu Ala Glu Val Leu Asp Pro Gln Asn Arg Glu Ser
65                  70                  75                  80
Arg Ser Pro Arg Arg Cys Val Arg Leu His Glu Ser Cys Leu Gly Gln
                85                  90                  95
Gln Val Pro Cys Cys Asp Pro Cys Ala Thr Cys Tyr Cys Arg Phe Phe
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Arg Pro Asp Gln Ala Leu Leu Pro Glu Leu Pro Gly Leu Gly Leu Arg
        35                  40                  45
Ala Pro Leu Lys Lys Thr Thr Ala Glu Gln Ala Glu Glu Asp Leu Leu
    50                  55                  60
Gln Glu Ala Gln Ala Leu Ala Glu Val Leu Asp Leu Gln Asp Arg Glu
65                  70                  75                  80
Pro Arg Ser Ser Arg Arg Cys Val Arg Leu His Glu Ser Cys Leu Gly
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Gln Gln Val Pro Cys Cys Asp Pro Cys Ala Thr Cys Tyr Cys Arg Phe
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Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp Met Ala Arg Tyr
        35                  40                  45
Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr Arg Gln Arg Tyr
    50                  55                  60
Gly Lys Arg Ser Ser Pro Glu Thr Leu Ile Ser Asp Leu Leu Met Lys
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ccagcggagg acatggccag atactactcg gcgctgcgac actacatcaa cctcatcacc 180
aggcagagat atggaaaacg atccagccca gagacactga tttcagacct cttgatgaga 240
gaaagcacag aaaatgttcc cagaactcgg cttgaagacc ctgcaatgtg gtgatgggaa 300
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<213> Homo sapien

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Ser Leu Leu Val Cys Leu Gly Ala Leu Ala Glu Ala Tyr Pro Ser Lys
            20   25   30
Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp Met Ala Arg Tyr
        35   40   45
Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr Arg Gln Arg Tyr
    50   55   60
Gly Lys Arg Ser Ser Pro Glu Thr Leu Ile Ser Asp Leu Leu Met Arg
65   70   75   80
Glu Ser Thr Glu Asn Val Pro Arg Thr Arg Leu Glu Asp Pro Ala Met
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<220>
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agttctgctg cagaaggcag aagctttggc ggaggtgcta gatccacaga accgcgagtc 240
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<220>
<223> Primer

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